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This Raw Listing contains the General Information
Section and up to first 5 pages.
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 1    <110> APPLICANT: McCarthy, Sean A.
 2    <120> TITLE OF INVENTION: NOVEL CRSP PROTEIN AND NUCLEIC ACID MOLECULES AND USES
 3          THEREFOR
 4    <130> FILE REFERENCE: MEI-008
 5    <140> CURRENT APPLICATION NUMBER: US/09/009,802B
 6    <141> CURRENT FILING DATE: 1998-01-20
 7    <150> EARLIER APPLICATION NUMBER: 08/842,898
 8    <151> EARLIER FILING DATE: 1997-04-17
 9    <150> EARLIER APPLICATION NUMBER: 60/071,589
10    <151> EARLIER FILING DATE: 1998-01-15
11    <160> NUMBER OF SEQ ID NOS: 19
12    <170> SOFTWARE: PatentIn Ver. 2.0
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14    <211> LENGTH: 2479
15    <212> TYPE: DNA
16    <213> ORGANISM: Homo sapiens
17    <220> FEATURE:
18    <221> NAME/KEY: CDS
19    <222> LOCATION: (38)..(1087)
20    <220> FEATURE:
21    <223> OTHER INFORMATION: 'n' at position 1146 may be any nucleotide
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26          acc ctg ctg tgc ctg ctg ctg gcg gcg gcg gtc ccc acg gcc ccc gcg  103
27          Thr Leu Leu Cys Leu Leu Leu Ala Ala Ala Val Pro Thr Ala Pro Ala
28                      10                  15                  20
29          ccc gct ccg acg gcg acc tcg gct cca gtc aag ccc ggc ccg gct ctc  151
30          Pro Ala Pro Thr Ala Thr Ser Ala Pro Val Lys Pro Gly Pro Ala Leu
31                  25                  30                  35
32          agc tac ccg cag gag gag gcc acc ctc aat gag atg ttc cgc gag gtt  199
33          Ser Tyr Pro Gln Glu Glu Ala Thr Leu Asn Glu Met Phe Arg Glu Val
34              40                  45                  50
35          gag gaa ctg atg gag gac acg cag cac aaa ttg cgc agc gcg gtg gaa  247
36          Glu Glu Leu Met Glu Asp Thr Gln His Lys Leu Arg Ser Ala Val Glu
37          55                  60                  65                  70
38          gag atg gag gca gaa gaa gct gct gct aaa gca tca tca gaa gtg aac  295
39          Glu Met Glu Ala Glu Glu Ala Ala Ala Lys Ala Ser Ser Glu Val Asn
40                          75                  80                  85
41          ctg gca aac tta cct ccc agc tat cac aat gag acc aac aca gac acg  343
42          Leu Ala Asn Leu Pro Pro Ser Tyr His Asn Glu Thr Asn Thr Asp Thr
43                  90                  95                  100
44          aac gtt gga aat aat acc atc cat gtg cac cga gaa att cac aag ata  391
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     47        acc aac aac cag act gga caa atg gtc ttt tca gag aca gtt atc aca     439
     48        Thr Asn Asn Gln Thr Gly Gln Met Val Phe Ser Glu Thr Val Ile Thr
     49                120                     125                     130
     50        tct gtg gga gac gaa gaa ggc aga agg agc cac gag tgc atc atc gac     487
     51        Ser Val Gly Asp Glu Glu Gly Arg Arg Ser His Glu Cys Ile Ile Asp
     52        135                     140                     145                150
     53        gag gac tgt ggg ccc agc atg tac tgc cag ttt gcc agc ttc cag tac     535
     54        Glu Asp Cys Gly Pro Ser Met Tyr Cys Gln Phe Ala Ser Phe Gln Tyr
     55                        155                     160                     165
     56        acc tgc cag cca tgc cgg ggc cag agg atg ctc tgc acc cgg gac agt     583
     57        Thr Cys Gln Pro Cys Arg Gly Gln Arg Met Leu Cys Thr Arg Asp Ser
     58                        170                     175                     180
     59        gag tgc tgt gga gac cag ctg tgt gtc tgg ggt cac tgc acc aaa atg     631
     60        Glu Cys Cys Gly Asp Gln Leu Cys Val Trp Gly His Cys Thr Lys Met
     61                        185                     190                     195
     62        gcc acc agg ggc agc aat ggg acc atc tgt gac aac cag agg gac tgc     679
     63        Ala Thr Arg Gly Ser Asn Gly Thr Ile Cys Asp Asn Gln Arg Asp Cys
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     65        cag ccg ggg ctg tgc tgt gcc ttc cag aga ggc ctg ctg ttc cct gtg     727
     66        Gln Pro Gly Leu Cys Cys Ala Phe Gln Arg Gly Leu Leu Phe Pro Val
     67        215                     220                     225                230
     68        tgc aca ccc ctg ccc gtg gag ggc gag ctt tgc cat gac ccc gcc agc     775
     69        Cys Thr Pro Leu Pro Val Glu Gly Glu Leu Cys His Asp Pro Ala Ser
     70                        235                     240                     245
     71        cgg ctt ctg gac ctc atc acc tgg gag cta gag cct gat gga gcc ttg     823
     72        Arg Leu Leu Asp Leu Ile Thr Trp Glu Leu Glu Pro Asp Gly Ala Leu
     73                        250                     255                     260
     74        gac cga tgc cct tgt gcc agt ggc ctc ctc tgc cag ccc cac agc cac     871
     75        Asp Arg Cys Pro Cys Ala Ser Gly Leu Leu Cys Gln Pro His Ser His
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     77        agc ctg gtg tat gtg tgc aag ccg acc ttc gtg ggg agc cgt gac caa     919
     78        Ser Leu Val Tyr Val Cys Lys Pro Thr Phe Val Gly Ser Arg Asp Gln
     79                280                     285                     290
     80        gat ggg gag atc ctg ctg ccc aga gag gtc ccc gat gag tat gaa gtt     967
     81        Asp Gly Glu Ile Leu Leu Pro Arg Glu Val Pro Asp Glu Tyr Glu Val
     82        295                     300                     305                310
     83        ggc agc ttc atg gag gag gtg cgc cag gag ctg gag gac ctg gag agg     1015
     84        Gly Ser Phe Met Glu Glu Val Arg Gln Glu Leu Glu Asp Leu Glu Arg
     85                        315                     320                     325
     86        agc ctg act gaa gag atg gcg ctg agg gag cct gcg gct gcc gcc gct     1063
     87        Ser Leu Thr Glu Glu Met Ala Leu Arg Glu Pro Ala Ala Ala Ala Ala
     88                        330                     335                     340
     89        gca ctg ctg gga agg gaa gag att tagatctgga ccaggctgtg ggtagatgtg    1117
     90        Ala Leu Leu Gly Arg Glu Glu Ile
     91                345                     350
W-->  92        caatagaaat agctaattta tttccccang tgtgtgcttt aagcgtgggc tgaccaggct    1177
     93        tcttcctaca tcttcttccc agtaagtttc ccctctggct tgacagcatg aggtgttgtg    1237
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105          gatttttctt ttgaggcatg cacatctgga attaaggtca aactaattct cacatccctc 1957
106          taaaagtaaa ctactgttag gaacagcagt gttctcacag tgtggggcag ccgtccttct 2017
107          aatgaagaca atgatattga cactgtccct ctttggcagt tgcattagta actttgaaag 2077
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110          ccacgtggag aaaatcaaac cgagcagggc tgtgtgaaac atggttgtaa tatgcgactg 2257
111          cgaacactga actctacgcc actccacaaa tgatgttttc aggtgtcatg gactgttgcc 2317
112          accatgtatt catccagagt tcttaaagtt taaagttgca catgattgta taagcatgct 2377
113          ttctttgagt tttaaattat gtataaacat aagttgcatt tagaaatcaa gcataaatca 2437
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123                          20                  25                  30
124          Lys Pro Gly Pro Ala Leu Ser Tyr Pro Gln Glu Glu Ala Thr Leu Asn
125                      35                  40                  45
126          Glu Met Phe Arg Glu Val Glu Glu Leu Met Glu Asp Thr Gln His Lys
127                  50                  55                  60
128          Leu Arg Ser Ala Val Glu Glu Met Glu Ala Glu Glu Ala Ala Ala Lys
129          65                  70                  75                  80
130          Ala Ser Ser Glu Val Asn Leu Ala Asn Leu Pro Pro Ser Tyr His Asn
131                          85                  90                  95
132          Glu Thr Asn Thr Asp Thr Asn Val Gly Asn Asn Thr Ile His Val His
133                      100                 105                 110
134          Arg Glu Ile His Lys Ile Thr Asn Asn Gln Thr Gly Gln Met Val Phe
135                  115                 120                 125
136          Ser Glu Thr Val Ile Thr Ser Val Gly Asp Glu Glu Gly Arg Arg Ser
137              130                 135                 140
138          His Glu Cys Ile Ile Asp Glu Asp Cys Gly Pro Ser Met Tyr Cys Gln
139          145                 150                 155                 160
140          Phe Ala Ser Phe Gln Tyr Thr Cys Gln Pro Cys Arg Gly Gln Arg Met
141                          165                 170                 175
142          Leu Cys Thr Arg Asp Ser Glu Cys Cys Gly Asp Gln Leu Cys Val Trp
143                          180                 185                 190
144          Gly His Cys Thr Lys Met Ala Thr Arg Gly Ser Asn Gly Thr Ile Cys
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146       Asp Asn Gln Arg Asp Cys Gln Pro Gly Leu Cys Cys Ala Phe Gln Arg
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148       Gly Leu Leu Phe Pro Val Cys Thr Pro Leu Pro Val Glu Gly Glu Leu
149       225              230              235              240
150       Cys His Asp Pro Ala Ser Arg Leu Leu Asp Leu Ile Thr Trp Glu Leu
151                       245              250              255
152       Glu Pro Asp Gly Ala Leu Asp Arg Cys Pro Cys Ala Ser Gly Leu Leu
153              260              265              270
154       Cys Gln Pro His Ser His Ser Leu Val Tyr Val Cys Lys Pro Thr Phe
155              275              280              285
156       Val Gly Ser Arg Asp Gln Asp Gly Glu Ile Leu Leu Pro Arg Glu Val
157              290              295              300
158       Pro Asp Glu Tyr Glu Val Gly Ser Phe Met Glu Glu Val Arg Gln Glu
159       305              310              315              320
160       Leu Glu Asp Leu Glu Arg Ser Leu Thr Glu Glu Met Ala Leu Arg Glu
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175       gtc ccc acg gcc ccc gcg ccc gct ccg acg gcg acc tcg gct cca gtc   96
176       Val Pro Thr Ala Pro Ala Pro Ala Pro Thr Ala Thr Ser Ala Pro Val
177                      20                  25                  30
178       aag ccc ggc ccg gct ctc agc tac ccg cag gag gag gcc acc ctc aat  144
179       Lys Pro Gly Pro Ala Leu Ser Tyr Pro Gln Glu Glu Ala Thr Leu Asn
180                      35                  40                  45
181       gag atg ttc cgc gag gtt gag gaa ctg atg gag gac acg cag cac aaa  192
182       Glu Met Phe Arg Glu Val Glu Glu Leu Met Glu Asp Thr Gln His Lys
183              50                  55                  60
184       ttg cgc agc gcg gtg gaa gag atg gag gca gaa gaa gct gct gct aaa  240
185       Leu Arg Ser Ala Val Glu Glu Met Glu Ala Glu Glu Ala Ala Ala Lys
186       65                  70                  75                  80
187       gca tca tca gaa gtg aac ctg gca aac tta cct ccc agc tat cac aat  288
188       Ala Ser Ser Glu Val Asn Leu Ala Asn Leu Pro Pro Ser Tyr His Asn
189                      85                  90                  95
190       gag acc aac aca gac acg aac gtt gga aat aat acc atc cat gtg cac  336
191       Glu Thr Asn Thr Asp Thr Asn Val Gly Asn Asn Thr Ile His Val His
192                      100                 105                 110
193       cga gaa att cac aag ata acc aac aac cag act gga caa atg gtc ttt  384
194       Arg Glu Ile His Lys Ile Thr Asn Asn Gln Thr Gly Gln Met Val Phe
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198                 130                    135                    140
199      cac gag tgc atc atc gac gag gac tgt ggg ccc agc atg tac tgc cag      480
200      His Glu Cys Ile Ile Asp Glu Asp Cys Gly Pro Ser Met Tyr Cys Gln
201      145                    150                    155                    160
202      ttt gcc agc ttc cag tac acc tgc cag cca tgc cgg ggc cag agg atg      528
203      Phe Ala Ser Phe Gln Tyr Thr Cys Gln Pro Cys Arg Gly Gln Arg Met
204                           165                    170                    175
205      ctc tgc acc cgg gac agt gag tgc tgt gga gac cag ctg tgt gtc tgg      576
206      Leu Cys Thr Arg Asp Ser Glu Cys Cys Gly Asp Gln Leu Cys Val Trp
207                 180                    185                    190
208      ggt cac tgc acc aaa atg gcc acc agg ggc agc aat ggg acc atc tgt      624
209      Gly His Cys Thr Lys Met Ala Thr Arg Gly Ser Asn Gly Thr Ile Cys
210                 195                    200                    205
211      gac aac cag agg gac tgc cag ccg ggg ctg tgc tgt gcc ttc cag aga      672
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215      Gly Leu Leu Phe Pro Val Cys Thr Pro Leu Pro Val Glu Gly Glu Leu
216      225                    230                    235                    240
217      tgc cat gac ccc gcc agc cgg ctt ctg gac ctc atc acc tgg gag cta      768
218      Cys His Asp Pro Ala Ser Arg Leu Leu Asp Leu Ile Thr Trp Glu Leu
219                           245                    250                    255
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221      Glu Pro Asp Gly Ala Leu Asp Arg Cys Pro Cys Ala Ser Gly Leu Leu
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223      tgc cag ccc cac agc cac agc ctg gtg tat gtg tgc aag ccg acc ttc      864
224      Cys Gln Pro His Ser His Ser Leu Val Tyr Val Cys Lys Pro Thr Phe
225                 275                    280                    285
226      gtg ggg agc cgt gac caa gat ggg gag atc ctg ctg ccc aga gag gtc      912
227      Val Gly Ser Arg Asp Gln Asp Gly Glu Ile Leu Leu Pro Arg Glu Val
228                 290                    295                    300
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231      305                    310                    315                    320
232      ctg gag gac ctg gag agg agc ctg act gaa gag atg gcg ctg agg gag     1008
233      Leu Glu Asp Leu Glu Arg Ser Leu Thr Glu Glu Met Ala Leu Arg Glu
234                           325                    330                    335
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